

The association of alanine aminotransferase within the normal and mildly elevated range with lipoproteins and apolipoproteins: the Insulin Resistance Atherosclerosis Study

C. Lorenzo · A. J. Hanley · M. J. Rewers · S. M. Haffner

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Abstract

Aims/hypothesis Markers of liver injury, such as alanine aminotransferase (ALT), have been associated with atherogenic lipoprotein changes. We examined the extent to which this association was explained by insulin resistance, adiposity, glucose tolerance and chronic inflammation.

Methods In this analysis we included 824 non-diabetic participants (age 40–69 years) in the Insulin Resistance Atherosclerosis Study. No participants reported excessive alcohol intake or treatment with lipid-lowering medications. Lipoproteins and apolipoproteins were measured by conventional methods and lipoprotein heterogeneity by nuclear magnetic resonance (NMR) spectroscopy.

Results ALT had a positive relationship with triacylglycerols, LDL-to-HDL-cholesterol ratio and apolipoprotein B

(ApoB) after adjusting for demographic variables ($p < 0.001$ for all three relationships). ALT was also associated with the following NMR lipoproteins: positively with large VLDL ($p < 0.001$), intermediate-density lipoprotein (IDL) ($p < 0.001$) and small LDL subclass particles ($p < 0.001$), and VLDL particle size ($p < 0.001$); and negatively with large LDL subclass particles ($p < 0.05$) and LDL ($p < 0.001$) and HDL particle sizes ($p < 0.01$). ALT remained associated with IDL and small LDL subclass particles and ApoB after adjusting for glucose tolerance, adiposity, directly measured insulin sensitivity and C-reactive protein.

Conclusions/interpretation ALT is associated with a wide range of atherogenic lipoprotein changes, which are partially explained by insulin resistance, adiposity, glucose tolerance and chronic inflammation. Because of the significant variability in the relationship between ALT and liver fat, further studies are needed to assess the extent of the lipoprotein changes using a direct measure of liver fat.

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Abbreviations

AIR	Acute insulin response
ALT	Alanine aminotransferase
Apo	Apolipoprotein
AST	Aspartate aminotransferase
FIB-4	Fibrosis 4 score
GGT	γ -Glutamyltransferase
HOMA-IR	Homeostasis model assessment of insulin resistance
IDF	International Diabetes Federation
IDL	Intermediate-density lipoprotein
IGT	Impaired glucose tolerance

IRAS	Insulin Resistance Atherosclerosis Study
NAFLD	Non-alcoholic fatty liver disease
NMR	Nuclear magnetic resonance
S_I	Insulin sensitivity index

Introduction

Chronic liver disease is a major cause of morbidity in the US population [1]. The epidemiology of chronic liver disease is rapidly changing. Viral and alcoholic hepatitis account for a significant number of cases, but the larger proportion is attributable to non-alcoholic fatty liver disease (NAFLD) (75.1% of all cases using data from the National Health and Nutrition Examination Survey 2005–2008) [2]. The clinical relevance of NAFLD is not limited to liver-related complications [3], but the relationship between NAFLD, cardiovascular disease and mortality is controversial [4–6]. This is so in part because of the differences between studies in terms of participants (elderly vs younger individuals), study design (cohort of individuals with NAFLD vs epidemiological studies) and criteria for exclusion (e.g. individuals who have elevated liver enzymes).

NAFLD may be considered a manifestation of the metabolic syndrome. Markers of liver injury including alanine aminotransferase (ALT) and γ -glutamyltransferase (GGT) have been shown to predict future diabetes [7, 8] and metabolic syndrome [9] independently of other risk factors. Measures of liver fat content and markers of liver injury have been associated with insulin resistance [10–13], overall and central adiposity [10–14], decreased adiponectin [10, 13], inflammation [14], and worsening insulin action on glucose and lipid metabolisms [10]. Measures of liver fat content and markers of liver injury have also been associated with dyslipidaemia including high triacylglycerols [10, 13, 15, 16], low HDL-cholesterol [10, 12–14, 16, 17], and increased VLDL particles [12], apolipoprotein B (ApoB) [12, 18] and apolipoprotein B-to-apolipoprotein A-1 ratio (ApoB-to-ApoA-1 ratio) [19]. However, none of these studies adjusted their results for insulin resistance, regional adiposity and glucose tolerance.

The conventional lipid panel may be insufficient to demonstrate the complete range of lipoprotein abnormalities in individuals with mildly elevated ALT. More sophisticated techniques such as nuclear magnetic resonance (NMR) spectroscopy [20] have been used to analyse lipoprotein heterogeneity in insulin resistance and adiposity [21]. Lipoprotein heterogeneity has been associated with carotid intima-media thickness [22], incident cardiovascular disease [23] and incident type 2 diabetes [24]. However, it is not known to what extent insulin resistance, adiposity, glucose tolerance and chronic inflammation explain the relationship between ALT, lipoproteins and apolipoproteins. Therefore, we examined lipoproteins, apolipoproteins and lipoprotein

heterogeneity as a function of ALT in non-diabetic participants in the Insulin Resistance Atherosclerosis Study (IRAS). ALT appears to be an appropriate marker for estimating liver fat in epidemiological studies. ALT has been shown to have a moderate to strong correlation with liver fat measured by magnetic resonance spectroscopy ($r=0.46$ in women and 0.62 in men) [14, 25].

Methods

Study population The IRAS is a large epidemiological study conducted at four clinical centres in the USA (Los Angeles, CA; Oakland, CA; San Antonio, TX; and the San Luis Valley, CO) to investigate the relationships between insulin resistance and cardiovascular disease. We enrolled 1,625 men and non-pregnant women from three ethnic groups (Hispanics, African-Americans and non-Hispanic whites) from 1992 through 1994. All participants provided written informed consent as approved by their respective centre's institutional review board.

This study includes data on non-diabetic participants in the IRAS ($n=824$). Reasons for participant exclusion in the analysis were the following: diabetes ($n=559$), treatment with lipid-lowering medications ($n=110$), excessive alcohol intake ($n=55$), missing information on ALT or lipoproteins ($n=71$), and ALT values three or more times above the normal range ($n=6$). Information on glucose tolerance was available for all participants, and information on insulin sensitivity index (S_I) for 792 of them.

Assessment of glucose tolerance and insulin sensitivity A full description of the study has been previously published [26]. Briefly, the IRAS protocol required two visits, 1 week apart, of approximately 4 h each at baseline. Participants were asked prior to each visit to fast for 12 h, to abstain from heavy exercise and alcohol for 24 h and to refrain from smoking on the morning of the examination. Age, sex, race/ethnicity, pharmacological treatment and alcohol intake were assessed by self-report. Anthropometric variables were measured using standard protocols. During the first visit, a 75 g oral glucose tolerance test was administered and blood was drawn immediately before and 2 h after the glucose load for repeat measurements of glucose and insulin concentrations. During the second baseline visit, insulin sensitivity and insulin secretion were determined using the frequently sampled intravenous glucose tolerance test with two modifications to the original protocol [27]. Insulin sensitivity, expressed as the S_I , was calculated using mathematical modelling methods (MINMOD version 3.0 [1994]; Los Angeles, CA, USA). Acute insulin response (AIR) was calculated as the mean of 2 min and 4 min insulin concentrations after glucose administration.

Laboratory procedures We used a glucose oxidase technique on an automated autoanalyser (Yellow Springs Instruments, Yellow Springs, OH, USA) to measure plasma glucose (interassay CV 3.2%) and a dextran-coated charcoal radioimmunoassay to determine insulin concentration (interassay CV 19%) [28]. The insulin assay displayed a high degree of cross-reactivity with proinsulin.

Total, LDL- and HDL-cholesterol were measured in plasma by the β -quantification procedure as described by the Lipid Research Clinics at the central IRAS laboratory (Medstar Laboratory, Washington, DC, USA) [29]. VLDL was isolated by preparative ultracentrifugation, and the VLDL (top) and bottom fractions were measured for cholesterol and triacylglycerol concentrations. HDL-cholesterol was measured after precipitation of ApoB-containing lipoproteins with $MnCl_2$ and heparin. The cholesterol content in the supernatant fraction was measured in a separate autoanalyser channel set to measure low cholesterol values. LDL-cholesterol was calculated as the difference between the HDL-cholesterol and the bottom cholesterol. Triacylglycerols were measured by enzymatic methods with the use of glycerol blanked assays on a Hitachi autoanalyser (Roche Molecular Biochemicals, Indianapolis, IN, USA) [29]. Interassay CV was 4% for LDL-cholesterol, HDL-cholesterol and triacylglycerols. Plasma total ApoB concentrations were assayed with an immunoprecipitation technique (SPQ kit from Instar Stillwater, MN, USA) and ApoA-I concentrations with a very sensitive ELISA assay at Medstar Laboratory [28]. Lower limits of detection for both ApoB and ApoA-I assays were 5 ng/ml and the interassay CV was 4.1%.

Lipoprotein subclass particle concentrations and average and VLDL, LDL and HDL particle diameters were measured by NMR spectroscopy (LipoScience, Raleigh, NC, USA) [20, 21]. In brief, the NMR signal of each subpopulation differs only slightly in frequency and line shape from the signals of neighbouring subpopulations. Measurement reproducibility of the individual signal amplitudes is inherently limited. To overcome this limitation, neighbouring subpopulations were grouped empirically into a smaller number of subclass categories (large, medium and small) so that the summed amplitudes of the individual subpopulation signals gave acceptable measurement precision (CV <10%). VLDL and LDL subclass particle concentrations are given in units of nanomoles per litre and those of HDL subclasses in micromoles per litre. Weighted-average VLDL, LDL and HDL particle sizes (in nanometres) were calculated as the sum of the diameter of each subclass multiplied by its relative mass percentage. The following subclasses were measured: large VLDL (including chylomicrons if present) (>60 nm), medium VLDL (35–60 nm), small VLDL (27–35 nm), intermediate-density lipoprotein (IDL) (23–27 nm), large LDL (21.2–23 nm), small LDL

(18–21.2 nm), large HDL (8.8–13 nm), medium HDL (8.2–8.8 nm) and small HDL (7.3–8.2 nm). CVs <4% were observed for total VLDL, LDL and HDL particle concentrations: <2% for VLDL size; <0.5% for LDL and HDL size; <10% for large, medium and small VLDL subclasses; <8% for large and small LDL subclasses; and <5% for large and small HDL subclasses. Higher CVs for IDL (<20%) and medium HDL (<35%) subclasses reflect their typically low concentrations [21].

ALT and aspartate aminotransferase (AST) were measured at the central IRAS laboratory with a Paramax PLA instrument (Baxter, Deerfield, IL, USA). ALT and AST were determined by enzymatic colorimetric assay using standard reagents by reaction rate assay based on the conversion of the reduced form of nicotinamide adenine dinucleotide to nicotinamide adenine dinucleotide (interassay CV <7%) [30]. One participant had an undetectable ALT concentration (range 0–1.47 μ kat/l), but none had an undetectable AST concentration (range 0.10–1.49 μ kat/l). C-reactive protein was measured by in-house ultrasensitive competitive immunoassay (antibodies and antigens from Calbiochem, La Jolla, CA, USA) with an interassay CV of 8.9% [31].

Variable definition We considered excessive alcohol consumption an intake of more than two drinks (28 g of alcohol) per day in men and more than one drink (14 g of alcohol) per day in women. Diabetes was defined as fasting glucose ≥ 7.0 mmol/l, 2 h plasma glucose ≥ 11.1 mmol/l and/or treatment with glucose-lowering medications. In individuals without diabetes, impaired glucose tolerance (IGT) was defined as 2 h plasma glucose between 7.8 and 11.0 mmol/l. We used BMI ≥ 30 kg/m² as the criterion for obesity and the definition of the metabolic syndrome by the 2005 International Diabetes Federation (IDF) to compute the NAFLD liver fat score [32]. Homeostasis model assessment of insulin resistance (HOMA-IR) was calculated as follows: HOMA-IR = fasting insulin (μ U/ml) \times fasting glucose (mmol/l)/22.5 [33]. We used ALT and the NAFLD liver fat score as surrogate markers of NAFLD [34], and NAFLD fibrosis score and Fibrosis 4 score (FIB-4) as markers of disease severity in NAFLD [35, 36]. NAFLD liver fat score = $-2.89 + (1.18 \times \text{IDF-defined metabolic syndrome [yes=1/no=0]}) + (0.45 \times \text{diabetes [yes=2/no=0]}) + (0.15 \times \text{fasting insulin [mU/l]}) + (0.04 \times \text{AST [U/l]}) - (0.94 \times \text{AST [U/l]/ALT [U/l]})$ [34]. An NAFLD score value ≥ -0.640 has been shown to predict NAFLD with a sensitivity of 86% and a specificity of 71%; a score ≥ 1.257 has a sensitivity of 59% and a specificity of 94% [34]. NAFLD fibrosis score = $-1.675 + (0.037 \times \text{age [years]}) + (0.094 \times \text{BMI [kg/m}^2\text{]}) + (1.13 \times \text{impaired fasting glucose/diabetes [yes=1/no=0]}) + (0.99 \times \text{AST/ALT ratio}) - (0.013 \times \text{platelet count [10}^9\text{/l]}) - (0.66 \times \text{albumin [g/dl]})$ [35]. FIB-4 = $\text{age [years]} \times \text{AST [U/l]} / (\text{platelet count [10}^9\text{/l]} \times (\text{ALT [U/l]})^{0.5})$ [36].

Statistical analysis Statistical analysis was carried out using SAS statistical software (version 9.2; SAS Institute, Cary, NC, USA). ALT has been shown to have stronger correlations with S_1 [13] and directly measured liver fat compared with AST [16]. We therefore used ALT as the marker of chronic liver injury.

We examined demographic and metabolic variables by quartiles of ALT using one-way analysis of covariance or logistic regression analyses. We assessed the strength of the relationship of ALT to lipoproteins, apolipoproteins and other metabolic variables by linear regression analysis. Log_e -transformed values of ALT, AST, triacylglycerols, fasting insulin, AIR, HOMA-IR and C-reactive protein were used to meet the assumptions of the test. Log_e transformation of (S_1+1) and (alcohol intake+1) were also used given that some participants had $S_1=0$ and alcohol intake=0 (5.6% and 46.8% of eligible participants, respectively). All probability values were two-sided. We considered a p value <0.05 as statistically significant.

Results

Among non-diabetic participants in the IRAS, rates of obesity, IGT and the metabolic syndrome were 28.9%, 30.1% and 40.1%, respectively. In addition, 38.9% and 16.9% of the participants had NAFLD liver fat scores ≥ -0.640 and ≥ 1.257 , respectively.

Table 1 presents the relationship between markers of liver fat, liver fibrosis, insulin sensitivity, insulin secretion,

adiposity and dyslipidaemia. The NAFLD liver fat score had a strong relationship with ALT, fasting insulin and S_1 ; a moderate relationship with BMI; and a weak relationship with AIR, triacylglycerols and HDL-cholesterol. NAFLD fibrosis score and FIB-4 had a negative relationship with ALT and were more weakly related to measures of insulin resistance, insulin secretion, adiposity and dyslipidaemia than was ALT. Correlation coefficients relating ALT to measures of insulin sensitivity, insulin secretion, adiposity and dyslipidaemia were lower than the corresponding coefficients of the NAFLD liver fat score. However, we chose ALT as the surrogate measure of liver fat, because the metabolic syndrome and insulin resistance were used to estimate the NAFLD liver fat score and thus could have had a confounding effect on the relationship between NAFLD liver fat score and dyslipidaemia.

Mean values of ALT and AST were 0.31 $\mu\text{kat/l}$ (range 0–1.47) and 0.38 $\mu\text{kat/l}$ (range 0.10–1.49), respectively. Compared with non-Hispanic whites, African-Americans had higher age- and sex-adjusted ALT (0.33 \pm 0.02 vs 0.27 \pm 0.01 $\mu\text{kat/l}$; $p<0.001$) and AST values (0.37 \pm 0.01 vs 0.34 \pm 0.01 $\mu\text{kat/l}$; $p=0.002$), but Hispanics had similar ALT (0.26 \pm 0.01 $\mu\text{kat/l}$; $p=0.784$) and AST values (0.35 \pm 0.01 $\mu\text{kat/l}$; $p=0.224$). Individuals with IGT had higher age-, sex- and race/ethnic-adjusted ALT values than did those with normal glucose tolerance (0.30 \pm 0.01 vs 0.27 \pm 0.01 $\mu\text{kat/l}$; $p=0.009$), but similar AST values (0.36 \pm 0.01 vs 0.35 \pm 0.01 $\mu\text{kat/l}$; $p=0.174$).

ALT was positively related to BMI, waist circumference, fasting and 2 h glucose, fasting insulin, HOMA-IR, C-reactive

Table 1 Spearman correlation coefficients between markers of NAFLD, liver fibrosis, insulin sensitivity, insulin secretion, adiposity and dyslipidaemia in non-diabetic participants in the IRAS

Marker	NAFLD liver fat score	NAFLD fibrosis score	FIB-4	Fasting insulin	S_1	AIR	BMI	Triacylglycerols	HDL-cholesterol
ALT	0.65***	-0.20***	-0.13***	0.35***	-0.29***	0.13***	0.18***	0.15***	-0.19***
NAFLD liver fat score ^a	–	0.05	-0.18***	0.85***	-0.66***	0.25***	0.56***	0.36***	-0.36***
NAFLD fibrosis score ^b		–	0.71***	0.15***	-0.19***	-0.10**	0.34***	0.11**	-0.08*
FIB-4 ^c			–	-0.12***	0.06	-0.06	-0.08*	0.04	0.03
Fasting insulin				–	-0.69***	0.35***	0.56***	0.34***	-0.32***
S_1					–	-0.31***	-0.55***	-0.30***	0.29***
AIR						–	0.22***	0.09*	-0.14***
BMI							–	0.21***	-0.19***
Triacylglycerols								–	-0.46***

^a NAFLD liver fat score as described by Kotronen et al [34]; NAFLD liver fat score = $-2.89 + (1.18 \times \text{IDF-defined metabolic syndrome}[\text{yes}=1/\text{no}=0]) + (0.45 \times \text{diabetes}[\text{yes}=2/\text{no}=0]) + (0.15 \times \text{fasting insulin} [\text{mU/l}]) + (0.04 \times \text{AST} [\text{U/l}]) - (0.94 \times \text{AST} [\text{U/l}]/\text{ALT} [\text{U/l}])$

^b NAFLD fibrosis score as described by Angulo et al [35]; NAFLD fibrosis score = $-1.675 + (0.037 \times \text{age}) + (0.094 \times \text{BMI} [\text{kg/m}^2]) + (1.13 \times \text{impaired fasting glucose/diabetes} [\text{yes}=1/\text{no}=0]) + (0.99 \times \text{AST} [\text{U/l}]/\text{ALT} [\text{U/l}]) - (0.013 \times \text{platelet count} [10^9/\text{l}]) - (0.66 \times \text{albumin} [\text{g/dl}])$

^c FIB-4 as described by Sterling et al [36]; FIB-4 = $\text{age} [\text{years}] \times \text{AST} [\text{U/l}] / (\text{platelet count} [10^9/\text{l}] \times (\text{ALT} [\text{U/l}])^{0.5})$

* $p<0.05$, ** $p<0.01$, *** $p<0.001$

Table 2 Metabolic variables by quartiles of ALT in 824 non-diabetic participants

Variable	Quartile				<i>p</i> for trend ^a
	1st	2nd	3rd	4th	
<i>N</i>	191	207	214	212	–
ALT, mean and range (μkat/l) ^b	0.13 (0, 0.17)	0.22 (0.18, 0.25)	0.31 (0.27, 0.37)	0.55 (0.38, 1.47)	–
AST, mean and range (μkat/l) ^b	0.29 (0.10, 0.70)	0.32 (0.13, 1.07)	0.37 (0.20, 0.92)	0.51 (0.22, 1.49)	–
Age (years) ^b	55.1±0.6	54.9±0.6	54.7±0.6	52.7±0.6	0.004
Female (%) ^b	79.6 (73.3, 84.7)	66.7 (60.0, 72.8)	50.5 (43.8, 57.1)	37.7 (31.5, 44.4)	<0.001
Ethnicity (%) ^b					
Non-Hispanic whites	39.8 (33.1, 46.9)	44.4 (37.8, 51.3)	36.0 (29.8, 42.6)	31.1 (25.3, 37.7)	0.021
African-Americans	22.0 (16.7, 28.4)	24.6 (19.2, 31.0)	28.5 (22.9, 34.9)	27.8 (22.2, 34.2)	0.124
Hispanics	38.2 (31.6, 45.3)	30.9 (25.0, 37.5)	35.5 (29.4, 42.2)	41.0 (34.6, 47.8)	0.351
BMI (kg/m ²)	27.4±0.4	27.9±0.4	28.7±0.4	29.9±0.4	<0.001
Waist circumference (cm)	87.3±0.9	89.0±0.9	91.4±0.8	93.6±0.9	<0.001
Fasting glucose (mmol/l)	5.29±0.04	5.33±0.04	5.43±0.04	5.57±0.04	<0.001
2 h glucose (mmol/l)	6.53±0.14	6.59±0.13	6.90±0.12	7.16±0.13	<0.001
Fasting insulin (pmol/l) ^c	59.4±2.4	68.4±3.0	81.0±3.0	105.6±4.2	<0.001
HOMA-IR ^c	2.32±0.11	2.68±0.12	3.24±0.14	4.35±0.24	<0.001
AIR (pmol/l) ^c	273.6±16.8	293.4±15.0	308.4±15.6	369.6±19.2	<0.001
S _I (×10 ⁻⁴ min ⁻¹ pmol ⁻¹) ^c	0.433±0.029	0.377±0.014	0.336±0.013	0.221±0.012	<0.001
Alcohol intake (g/day) ^c	1.35±0.17	1.34±0.15	1.14±0.14	1.41±0.16	0.995
C-reactive protein (nmol/l) ^c	13.9±1.1	17.9±1.5	18.1±1.3	20.2±1.7	0.003
Metabolic syndrome (%)	23.2 (17.5, 30.1)	33.6 (27.3, 40.7)	45.3 (38.5, 52.2)	54.7 (47.3, 61.9)	<0.001
NAFLD liver fat score	-2.71±0.20	-1.02±0.19	-0.02±0.18	1.20±0.19	<0.001
NAFLD liver fat score≥-0.640 (%)	10.0 (6.5, 15.3)	32.4 (26.2, 39.4)	49.3 (42.5, 56.2)	78.0 (71.4, 83.3)	<0.001
NAFLD fibrosis score	0.01±0.12	-1.17±0.11	-1.30±0.11	-1.21±0.11	<0.001
FIB-4	1.43±0.04	1.15±0.04	1.14±0.04	1.25±0.04	0.006
Lipoproteins (mmol/l)					
Total cholesterol	5.25±0.08	5.49±0.08	5.45±0.07	5.51±0.08	0.060
LDL-cholesterol	3.48±0.07	3.70±0.06	3.63±0.06	3.72±0.06	0.041
HDL-cholesterol	1.22±0.03	1.23±0.02	1.17±0.02	1.14±0.02	0.010
LDL-to-HDL-cholesterol ratio	3.09±0.10	3.31±0.09	3.41±0.09	3.61±0.09	<0.001
Triacylglycerols ^c	1.16±0.05	1.26±0.05	1.31±0.05	1.43±0.06	<0.001
Apolipoproteins (g/l)					
ApoA-1	1.27±0.02	1.31±0.02	1.30±0.02	1.29±0.02	0.639
ApoB	0.98±0.02	1.02±0.02	1.05±0.02	1.12±0.02	<0.001
ApoB-to-ApoA-1 ratio	0.81±0.02	0.81±0.02	0.84±0.02	0.91±0.02	0.001
NMR measures					
Total VLDL particles (nmol/l)	60.7±2.2	64.2±2.1	64.7±2.0	68.4±2.1	0.023
Large	2.30±0.21	2.91±0.19	2.93±0.19	3.34±0.20	0.001
Medium	17.0±0.9	17.9±0.9	17.9±0.9	18.2±0.9	0.670
Small	41.4±1.5	43.4±1.4	43.9±1.3	46.9±1.4	0.015
Total LDL particles (nmol/l)	1,077±29	1,152±26	1,164±26	1,273±27	<0.001
IDL	37.1±1.9	42.5±1.7	42.5±1.7	47.8±1.8	<0.001
Large	524±16	527±14	486±14	465±15	0.002
Small	516±34	583±31	637±31	760±32	<0.001
Total HDL particles (μmol/l)	30.7±0.4	31.5±0.3	31.6±0.3	30.8±0.4	0.879
Large	4.96±0.18	5.22±0.17	4.74±0.16	4.39±0.17	0.007
Medium	2.38±0.25	2.60±0.23	2.92±0.22	2.57±0.24	0.446
Small	23.4±0.4	23.7±0.3	23.9±0.3	23.9±0.4	0.341

Table 2 (continued)

Variable	Quartile				<i>p</i> for trend ^a
	1st	2nd	3rd	4th	
Size (nm)					
VLDL	47.6±0.8	47.8±0.7	49.4±0.7	50.8±0.7	0.001
LDL	21.55±0.06	21.42±0.05	21.31±0.05	21.12±0.05	<0.001
HDL	9.01±0.03	8.98±0.03	8.92±0.03	8.84±0.03	<0.001

Data are *n*, means ± SE, % (95% CI), or mean and range

^a Results adjusted for age, sex, race/ethnicity and clinic

^b Non-adjusted results

^c Log_e-transformed variables

protein, NAFLD liver fat score, LDL-cholesterol, LDL-to-HDL-cholesterol ratio, triacylglycerols, ApoB and ApoB-to-ApoA-1 ratio after adjusting for age, sex, race/ethnicity and clinic (Table 2). ALT was negatively related to age, S₁, HDL-cholesterol, NAFLD fibrosis score and FIB-4. ALT had no significant relationship with alcohol intake (in the truncated distribution of the intake), total cholesterol or ApoA-1. NMR spectroscopy uncovered additional associations for ALT: positive relationships with total VLDL particles, large and small

VLDL subclass particles, total LDL particles, IDL and small LDL subclass particles, and VLDL particle size; and inverse relationships with large LDL and large HDL subclass particles, and LDL and HDL particle sizes. Figure 1 presents the relative magnitude of the change in lipoproteins and apolipoproteins across ALT categories.

We fitted a multiple linear regression model to data with ALT as the dependent variable and age, sex, race/ethnicity, clinic, IGT, alcohol intake and waist circumference as

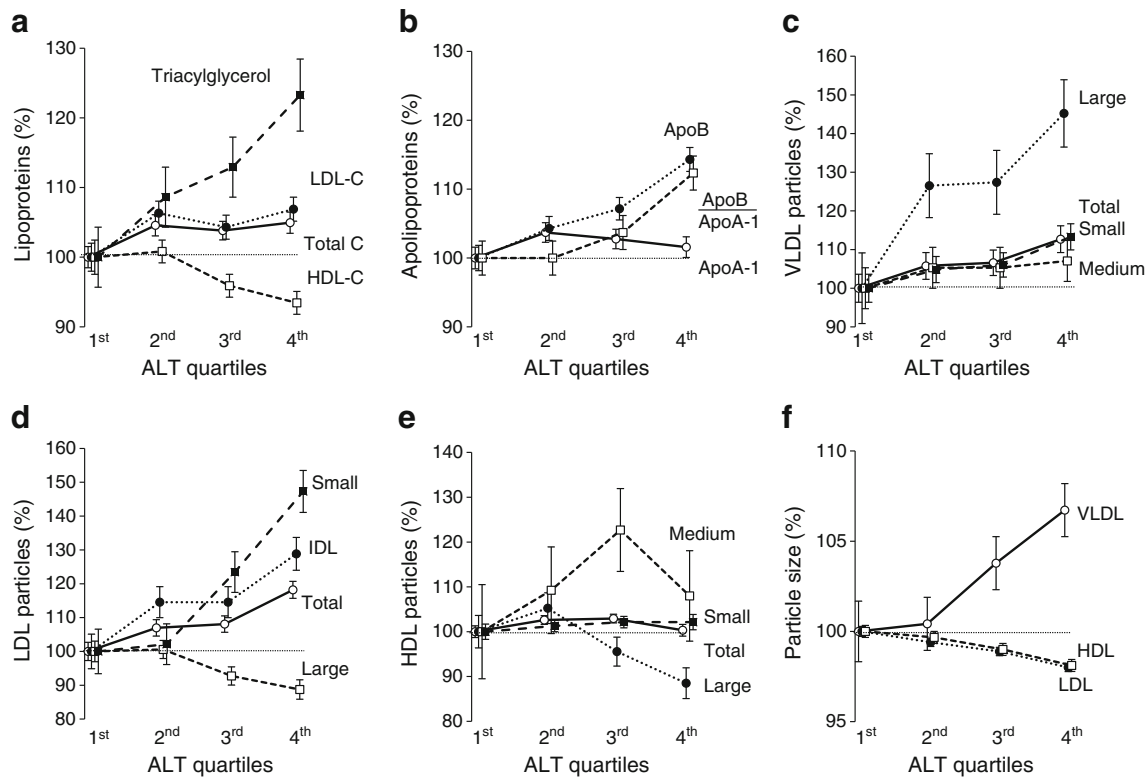


Fig. 1 (a–f) Relative change of lipoproteins, apolipoproteins and lipoprotein composition across ALT quartiles. All results adjusted for age, sex, race/ethnic origin and clinic. Range of ALT within each quartile was as follows: first quartile 0–0.17 μ kat/l; second quartile

0.18–0.25 μ kat/l; third quartile 0.26–0.37 μ kat/l; fourth quartile 0.38–1.47 μ kat/l. The first quartile (ALT 0–0.17 μ kat/l) was used as the referent category. C, cholesterol

Table 3 Multiple linear regression analysis with log ALT as the dependent variable

Independent variable	Model 1 $\beta \pm SE$	Model 2 $\beta \pm SE$	Model 3 $\beta \pm SE$
Intercept	3.482±0.470***	3.813±0.516***	4.400±0.483***
Log _e age ^a	-0.047±0.018**	-0.047±0.018**	-0.057±0.018**
Women vs men	-0.314±0.040***	-0.319±0.040***	-0.336±0.038***
Race/ethnicity			
African-Americans vs non-Hispanic whites	0.042±0.053	0.002±0.054	0.000±0.053
Hispanics vs non-Hispanic whites	0.090±0.049	0.075±0.049	0.065±0.048
Log _e alcohol intake ^a	0.001±0.019	0.008±0.019	0.019±0.019
IGT vs normal glucose tolerance	0.040±0.040	0.035±0.043	0.003±0.043
Waist circumference ^a	0.088±0.019***	0.036±0.022	–
Log _e S ₁ ^a	–	-0.097±0.023***	-0.117±0.023***
Log _e AIR ^a	–	0.036±0.020	0.036±0.019
Metabolic syndrome (yes vs no)	–	–	0.122±0.041**
Log _e C-reactive protein ^a	–	–	-0.005±0.020
R ² for the model	22.5%	24.6%	27.2%

Results also adjusted for clinic

^a $\beta \pm SE$ expressed per 1 SD unit increase

** $p < 0.01$, *** $p < 0.001$

independent variables (Table 3). Age, sex and waist circumference were the only variables that had independent associations with ALT. Because of the close relationship between central adiposity, insulin sensitivity and liver fat, we added S₁ and AIR as independent variables in a second model. S₁ explained the relationship between waist circumference and ALT. Finally, the relationship of metabolic syndrome and C-reactive protein to ALT was assessed in a third model. Both S₁ and metabolic syndrome had an independent relationship with ALT. This model explained 27.2% of the ALT variance.

We also fitted additional multiple linear regression models to data with individual lipoproteins and apolipoproteins as the dependent variable and age, sex, race/ethnicity, clinic, alcohol intake and ALT as independent variables (Table 4). ALT was directly associated with total and LDL-cholesterol, LDL-to-HDL-cholesterol ratio, triacylglycerols, ApoB and ApoB-to-ApoA-1 ratio. ALT was also related to the following NMR lipoproteins: directly to large and small VLDL subclass particles, total LDL particles, IDL and small LDL subclass particles, and VLDL particle size; and inversely to large LDL subclass particles and LDL and HDL particle sizes (model 1). However, only the relationship of ALT to IDL and small LDL subclass particles and ApoB remained significant after adjusting for glucose tolerance, waist circumference and S₁ (model 2). The additional adjustment for C-reactive protein had no significant impact on these relationships (model 3). In models that had the metabolic syndrome as a covariate, ALT was associated with IDL and small LDL subclass particles, VLDL and LDL particle sizes, and ApoB (model 4). The relationship between ALT and IDL subclass particles and ApoB remained statistically significant even after the additional adjustment for waist

circumference, glucose tolerance, S₁ and C-reactive protein (model 5).

In multiple linear regression models with individual lipoproteins and apolipoproteins as the dependent variable and age, sex, race/ethnicity, clinic, alcohol intake and ALT as independent variables, we assessed heterogeneity by introducing interaction terms sex×ALT, or ethnicity×ALT. We detected no p value reaching statistical significance for interaction terms sex×ALT. This indicates that sex had no significant effect on the relationship between ALT and each individual lipoprotein and apolipoprotein. However, ethnicity had a significant interaction effect on the relationship of ALT to triacylglycerol ($p=0.022$) and total VLDL particles ($p=0.043$). Figure 2 presents the relationship of ALT to S₁, ApoB, total LDL particles and small LDL subclass particles by sex, race/ethnicity and obesity (BMI <30 and ≥30 kg/m²). ALT had a statistically significant relationship with S₁ and ApoB in all categories and with total LDL particles and small LDL subclass particles in all but non-Hispanic whites.

Discussion

In non-diabetic individuals, ALT has direct relationships with triacylglycerols, LDL-to-HDL-cholesterol ratio, ApoB, and ApoB-to-ApoA-1 ratio after adjusting for age, sex, race/ethnicity, clinic and alcohol intake. Analysis of lipoprotein heterogeneity by NMR spectroscopy reveals additional associations: (1) ALT is directly related to large and small VLDL subclass particles, total LDL particles, IDL and small LDL subclass particles, and VLDL particle size; and (2) ALT is inversely related to large LDL and large HDL

Table 4 Relationship between log ALT as the independent variable and individual lipoproteins and apolipoproteins as the dependent variable

Dependent variable	Model 1 ^a β (95% CI)	Model 2 ^b β (95% CI)	Model 3 ^c β (95% CI)	Model 4 ^d β (95% CI)	Model 5 ^e β (95% CI)
Lipoproteins					
Total cholesterol	0.95 (0.11, 1.78)*	0.84 (−0.06, 1.74)	0.86 (−0.05, 1.77)	0.77 (−0.09, 1.62)	0.82 (−0.09, 1.73)
LDL-cholesterol	0.80 (0.13, 1.46)*	0.65 (−0.06, 1.37)	0.74 (0.02, 1.48)*	0.67 (−0.01, 1.34)	0.72 (−0.01, 1.44)
HDL-cholesterol	−0.23 (−0.48, 0.02)	0.12 (−0.13, 0.38)	0.08 (−0.18, 0.34)	0.06 (−0.18, 0.29)	0.16 (−0.09, 0.40)
LDL-to-HDL ratio	0.17 (0.07, 0.27)***	0.07 (−0.03, 0.18)	0.09 (−0.02, 0.19)	0.08 (−0.01, 0.17)	0.06 (−0.04, 0.16)
Log triacylglycerols	0.07 (0.03, 0.11)***	0.02 (−0.02, 0.06)	0.02 (−0.02, 0.06)	0.02 (−0.01, 0.06)	0.00 (−0.03, 0.04)
Apolipoproteins					
ApoA-1	1.17 (−0.80, 3.13)	2.19 (0.11, 4.27)*	1.83 (−0.27, 3.93)	2.09 (0.11, 4.07)*	2.01 (−0.08, 4.10)
ApoB	4.67 (2.84, 6.49)***	3.70 (1.79, 5.61)***	3.82 (1.89, 5.76)***	3.21 (1.43, 5.00)***	3.37 (1.49, 5.26)***
ApoB-to-ApoA-1 ratio	0.27 (0.06, 0.48)*	0.14 (−0.07, 0.36)	0.17 (−0.06, 0.39)	0.10 (−0.10, 0.31)	0.12 (−0.10, 0.34)
NMR particles					
Total VLDL	2.15 (−0.09, 4.39)	1.29 (−1.09, 3.66)	1.14 (−1.28, 3.56)	0.56 (−1.65, 2.78)	0.60 (−1.75, 2.95)
Large	0.37 (0.15, 0.58)***	0.15 (−0.07, 0.37)	0.13 (−0.09, 0.36)	0.17 (−0.03, 0.38)	0.08 (−0.14, 0.30)
Medium	0.03 (−0.92, 0.98)	−0.26 (−1.28, 0.76)	−0.34 (−1.38, 0.70)	−0.62 (−1.57, 0.33)	−0.59 (−1.59, 0.42)
Small	1.75 (0.25, 3.26)*	1.40 (−0.19, 2.98)	1.35 (−0.26, 2.96)	1.01 (−0.49, 2.51)	1.10 (−0.48, 2.68)
Total LDL	57.8 (29.0, 86.6)***	28.0 (−1.74, 57.8)	28.4 (−1.83, 58.7)	28.0 (0.30, 55.7)*	19.5 (−9.79, 48.8)
IDL	3.44 (1.54, 5.33)***	2.65 (0.64, 4.65)**	2.61 (0.57, 4.65)*	2.52 (0.61, 4.43)**	2.37 (0.33, 4.41)*
Large	−18.9 (−34.6, −3.19)*	−9.90 (−26.6, 6.80)	−10.2 (−27.2, 6.74)	−9.82 (−25.6, 5.93)	−7.78 (−24.6, 9.00)
Small	73.2 (39.0, 107.5)***	35.3 (0.00, 70.5)*	36.0 (0.08, 72.0)*	35.3 (2.74, 67.8)*	24.9 (−9.63, 59.4)
Total HDL	−0.01 (−0.39, 0.37)	0.02 (−0.39, 0.42)	0.05 (−0.37, 0.46)	0.06 (−0.33, 0.44)	0.06 (−0.35, 0.47)
Large	−0.16 (−0.34, 0.02)	0.05 (−0.14, 0.23)	0.02 (−0.17, 0.20)	0.01 (−0.16, 0.19)	0.06 (−0.12, 0.24)
Medium	0.16 (−0.09, 0.41)	0.12 (−0.15, 0.39)	0.12 (−0.16, 0.39)	0.12 (−0.14, 0.37)	0.11 (−0.16, 0.39)
Small	−0.01 (−0.39, 0.37)	−0.15 (−0.56, 0.25)	−0.09 (−0.49, 0.32)	−0.07 (−0.46, 0.31)	−0.11 (−0.52, 0.29)
NMR particle size					
VLDL	1.30 (0.54, 2.07)***	0.48 (−0.31, 1.27)	0.41 (−0.39, 1.22)	0.92 (0.15, 1.70)*	0.36 (−0.45, 1.17)
LDL	−0.12 (−0.18, −0.07)***	−0.05 (−0.11, 0.01)	−0.06 (−0.12, 0.005)	−0.06 (−0.12, −0.01)*	−0.04 (−0.10, 0.02)
HDL	−0.05 (−0.08, −0.02)**	−0.01 (−0.04, 0.02)	−0.02 (−0.05, 0.02)	−0.01 (−0.04, 0.02)	−0.01 (−0.04, 0.02)

β ± SE expressed per 1 SD unit increase in log ALT

Adjustment models:

^a Model 1: Results adjusted for log age, sex, race/ethnicity, clinic and log alcohol intake

^b Model 2: Results adjusted for variables in model 1 plus IGT, waist circumference and S_1

^c Model 3: Results adjusted for variables in model 2 plus C-reactive protein

^d Model 4: Results adjusted for variables in model 1 plus metabolic syndrome

^e Model 5: Results adjusted for variables in model 4 plus IGT, waist circumference, S_1 and C-reactive protein

* $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$

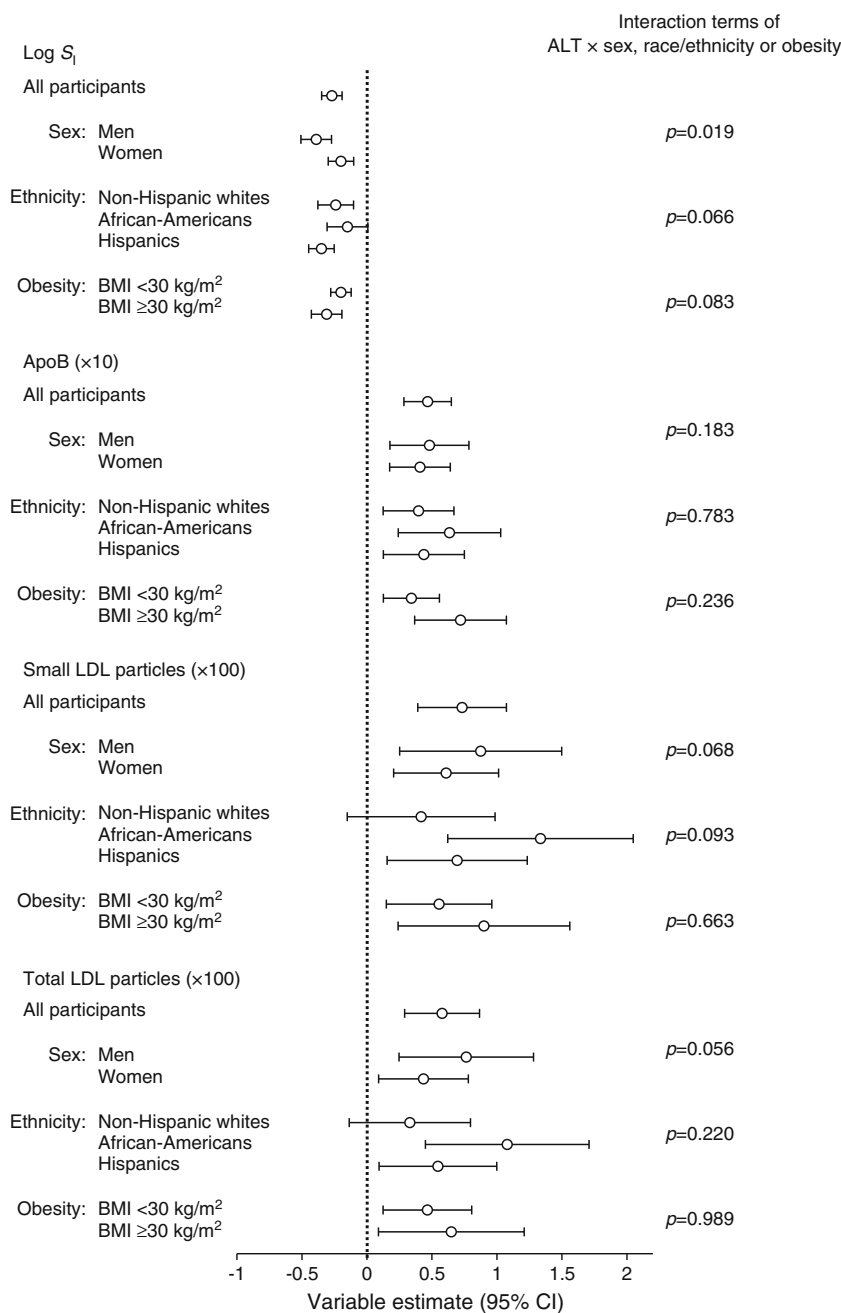
subclass particles and LDL and HDL particle sizes. The relationship of ALT to ApoB and IDL and small LDL subclass particles is independent of the effect of insulin sensitivity, central adiposity and glucose tolerance.

NAFLD is a feature of the metabolic syndrome and denotes hepatic insulin resistance [10, 14, 37]. Liver fat content measured by proton magnetic resonance spectroscopy partially explains the variation in triacylglycerols, HDL-cholesterol and insulin concentrations independently of the effect of intra-abdominal and overall adiposity [14].

Individuals with NAFLD have increased cholesterol synthesis [38] and overproduction of large VLDL subclass particles and ApoB [12, 39]. Overproduction of VLDL particles results in the generation of small, dense LDL subclass particles and low HDL-cholesterol, all of which are characteristic traits of individuals with type 2 diabetes [12].

Multiple studies have described significant relationships between liver fat content, or markers of liver injury such as ALT and GGT, and lipoproteins (triacylglycerols, HDL-

Fig. 2 Heterogeneity analyses on the relationship of ALT (independent variable) to S_1 , ApoB, small LDL and total LDL (dependent variable). Age, sex, race/ethnicity, clinic and alcohol consumption were also included as covariates in all models. Heterogeneity was assessed by introducing interaction terms sex \times ALT, ethnicity \times ALT, or obesity \times ALT



cholesterol and large VLDL particles) [10, 12–14, 16, 17] and apolipoproteins (ApoB and ApoB-to-ApoA-1 ratio [12, 18, 19]. However, few studies have adjusted their results for regional adiposity, glucose tolerance or insulin resistance. Since ALT and insulin resistance are both related to dyslipidaemia, an unresolved issue is to establish the independent effect of liver fat on lipoproteins and apolipoproteins. ALT and GGT have been associated with triacylglycerols after adjusting for fasting insulin, glucose tolerance status and adiposity [40]. ALT has also been associated with high triacylglycerol and/or low HDL-cholesterol concentrations after taking into account the effect of the other metabolic syndrome-related

disorders [41]. Lifestyle intervention and treatment with rimonabant have also been shown to reduce liver fat content, adiposity and dyslipidaemia in individuals with and without diabetes [42, 43].

Our present study is in agreement with previous reports that have used conventional methods to measure dyslipidaemic abnormalities following liver fat accumulation. Specifically, ALT has been associated with triacylglycerols, LDL-to-HDL-cholesterol ratio, ApoB, and ApoB-to-ApoA-1 ratio in non-diabetic individuals. However, our study has novel findings: (1) the relationships of ALT with triacylglycerols, LDL-to-HDL-cholesterol ratio and ApoB are not

fully explained by glucose tolerance and central adiposity; and (2) the association between ALT and ApoB is not explained by a direct measure of insulin sensitivity. ApoB reflects the total number of atherogenic particles (VLDL, IDL and LDL subclass particles), but LDL particles usually contribute to more than 90% of the total amount of ApoB [44]. The production of ApoB is increased in individuals with NAFLD [12]. Our study differs somewhat from previous reports by disclosing a weak relationship between ALT and HDL-cholesterol (see the non-linear relationship by quartiles in Table 2). The absence of a relationship between ALT and ApoA-1, the major apolipoprotein of HDL particles, suggests that liver fat may have a small influence on reverse cholesterol transport and protection against the atherosclerotic process (both of which are key functions of HDL-cholesterol particles and ApoA-1) [45].

Sophisticated techniques such as NMR spectroscopy have been helpful in assessing lipoprotein heterogeneity in insulin-resistant states including obesity [20]. Weight loss induced by rimonabant has been associated with a reduction in liver fat and small LDL subclass particles and an increase in large LDL subclass particles and LDL and HDL particle sizes [43]. Our results further expand the extent of lipoprotein changes using ALT as a surrogate marker of liver injury. ALT is directly related to large and small VLDL subclass particles, VLDL particle size, total LDL particles, and IDL and small LDL subclass particles. ALT is inversely related to large LDL subclass particles and LDL and HDL particle sizes. Insulin resistance, central adiposity and liver fat appear to have distinct effects on the relationship of ALT to lipoprotein composition [46]. These relationships are partially explained by insulin resistance, central adiposity and glucose tolerance except for those involving IDL and small LDL subclass particles. Thus, high IDL and small LDL subclass particles as well as high ApoB may be characteristics of individuals with mildly elevated ALT.

Intra-abdominal fat may account for ethnic differences in liver fat [47]. African-Americans tend to have less intra-abdominal fat, liver fat and insulin sensitivity compared with non-Hispanic whites. In our study, African-Americans had higher ALT levels in comparison with non-Hispanic whites and Hispanics. This is probably due to the lower insulin sensitivity in African-Americans compared with the other two populations. Central adiposity and insulin sensitivity largely explain ethnic differences in ALT.

ALT is an imperfect measure of NAFLD and not one that may be suitable for clinical decision making. A better measure is desirable, but it is not available in the IRAS. However, ALT has been shown to correlate well with directly measured liver fat by magnetic resonance spectroscopy ($r=0.46$ in women and 0.62 in men [14], making ALT appropriate for estimating liver fat in epidemiological studies [25]. The IRAS and other studies have shown that ALT

correlates with direct measures of insulin resistance and sensitivity and measures of dyslipidaemia [11, 48] to predict future development of the metabolic syndrome [9] and type 2 diabetes [7, 48]. In search of a better measure, we have examined the NAFLD liver fat score (a measure of liver fat) and the NAFLD fibrosis score and FIB-4 (both indices of fibrosis in NAFLD). An NAFLD liver fat score of ≥ -0.640 is considered optimal for detecting individuals with NAFLD and is quite prevalent in non-diabetic participants in the IRAS. However, the NAFLD liver fat score may not be the most suitable surrogate measure to analyse the relationship between NAFLD and dyslipidaemia. The NAFLD liver fat score formula includes both fasting insulin (a measure of insulin resistance) and the metabolic syndrome (a clustering of risk factors including triacylglycerol and HDL-cholesterol levels). Therefore, the relationship between NAFLD liver fat score and dyslipidaemia may be driven by the relationship that lipoproteins have with insulin resistance and the metabolic syndrome.

Similarly, surrogate indices of fibrosis in individuals with NAFLD do not appear appropriate for studying the relationship between NAFLD and dyslipidaemia. The NAFLD fibrosis score and FIB-4 are inversely related to ALT and more weakly associated with insulin sensitivity and levels of triacylglycerol and HDL-cholesterol than is ALT. In individuals with biopsy-proven NAFLD, high Kleiner scores (the gold standard to assess disease severity in NAFLD) are associated with lower cholesterol and triacylglycerol concentrations compared with low Kleiner scores [49]. Surrogate markers of liver fibrosis may indicate severity of the disease in NAFLD, but the development of liver fibrosis may have consequences for lipoprotein concentration and composition beyond those of liver fat.

Major strengths of our study include the use of data from a large, well-characterised multiethnic cohort of US adults, assessment of glucose tolerance, and measurement of insulin sensitivity by a direct method and lipoprotein heterogeneity by NMR spectroscopy. Our study has also several limitations. It lacks information on directly measured liver fat and GGT and has no information on serology for hepatitis B and C or severity of NAFLD. However, our results also indicate that the relationship of ALT with lipoprotein and apolipoprotein abnormalities is consistent across sex and race/ethnic categories.

In summary, NMR spectroscopy expands the range of atherogenic lipoprotein changes that conventional methods detect in individuals with mildly elevated ALT. The relationship between ALT and lipoprotein composition is partially explained by insulin resistance, adiposity and glucose tolerance status. Because of the significant ALT variability with changes in liver fat, studies with data on directly measured liver fat are needed to assess the extent of the lipoprotein changes in NAFLD.

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